

Figure S5

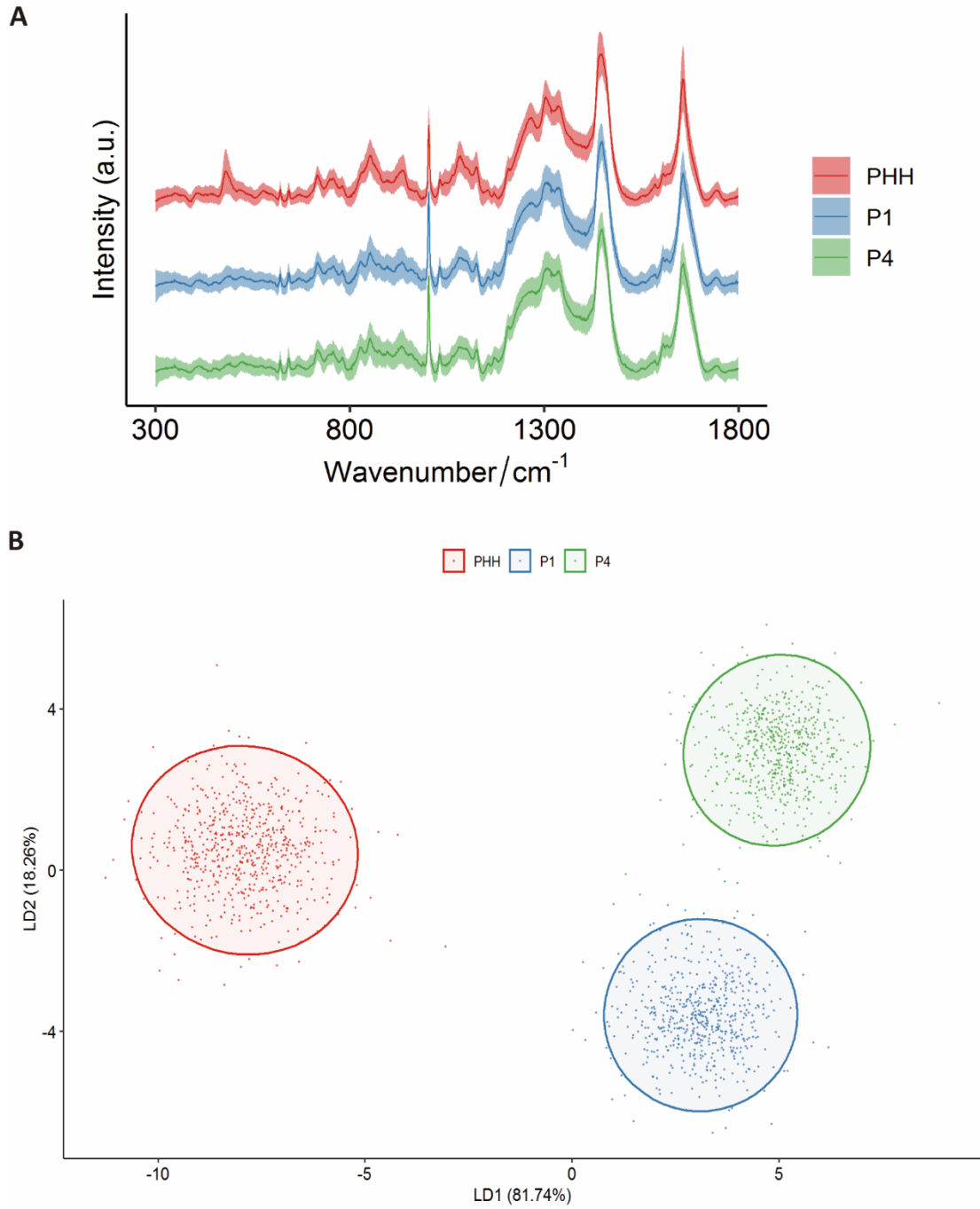


Figure S5. Raman spectroscopy and classification analysis for PHH (Lot:201678901), ProliHHs P1 and P4. (A) The averaged spectra ($n = 1829$) collected by PHH ($n = 619$), P1 ($n = 595$) and P4 ($n = 615$) on fingerprint region. (B) Linear discriminant analysis clearly distinguished three cell groups. (The red, blue, and green colors represent PHH, ProliHHs P1 and P4 cells, respectively. PHH: primary human hepatocytes, ProliHHs: proliferating human hepatocytes, P1: passage 1, P4: passage 4)

Figure S6

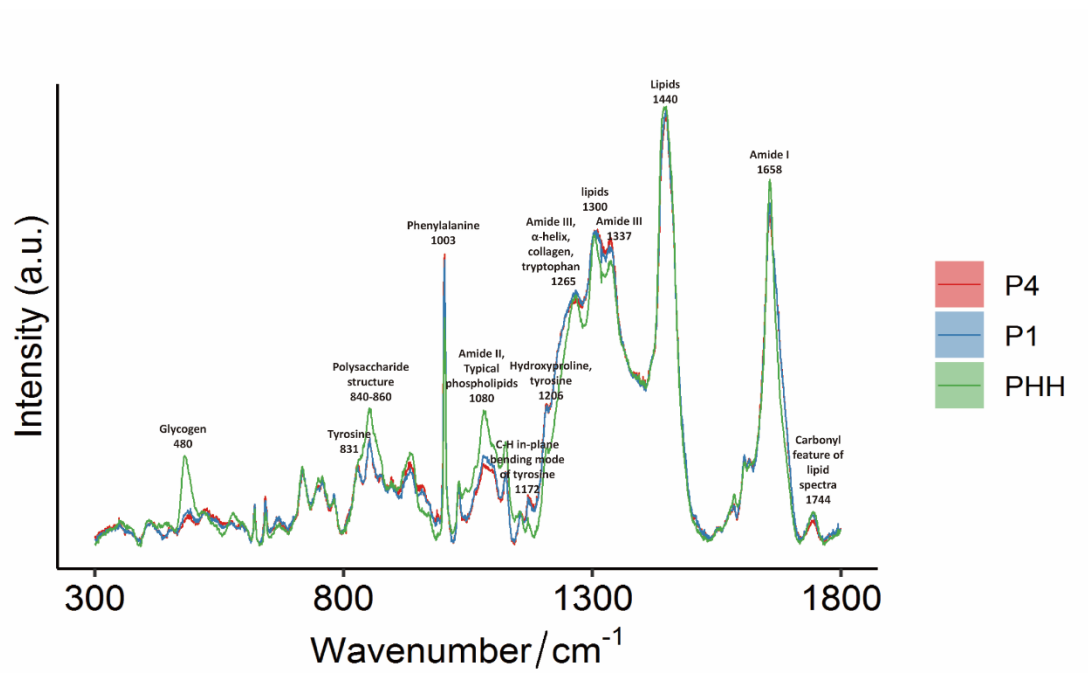


Figure S6. The biochemical molecules represented by the specific Raman bands in the average spectral (Lot:201678901).

Figure S7

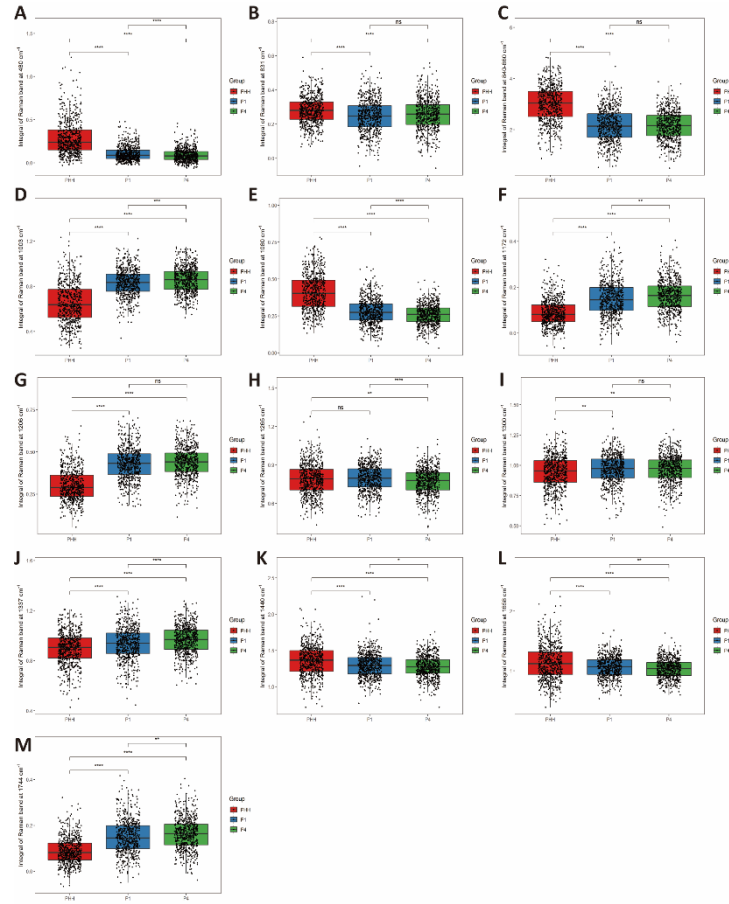


Figure S7. The peak area were semi-quantitative to compare differences of the specific Raman bands (A) 480 cm^{-1} (glycogen), (B) 831 cm^{-1} (tyrosine), (C) $840\text{--}860\text{ cm}^{-1}$ (polysaccharide structure), (D) 1003 cm^{-1} (phenylalanine), (E) 1080 cm^{-1} (amide II, typical phospholipid), (F) 1172 cm^{-1} (C-H in-plane bending mode of tyrosine), (G) 1206 cm^{-1} (hydroxyproline, tyrosine), (H) 1265 cm^{-1} (α -helix, collagen, tryptophan), (I) 1300 cm^{-1} (lipids), (J) 1337 cm^{-1} (amide III), (K) 1440 cm^{-1} (lipids), (L) 1658 cm^{-1} (amide I), (M) 1744 cm^{-1} (carbonyl feature of lipid spectra) in PHH (Lot:201678901), ProliHHs P1 and P4. The results represent median, ns $P \geq 0.05$, * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, **** $P < 0.0001$. (PHH: primary human hepatocytes, ProliHHs: proliferating human hepatocytes, P1: passage 1, P4: passage 4)

Figure S8

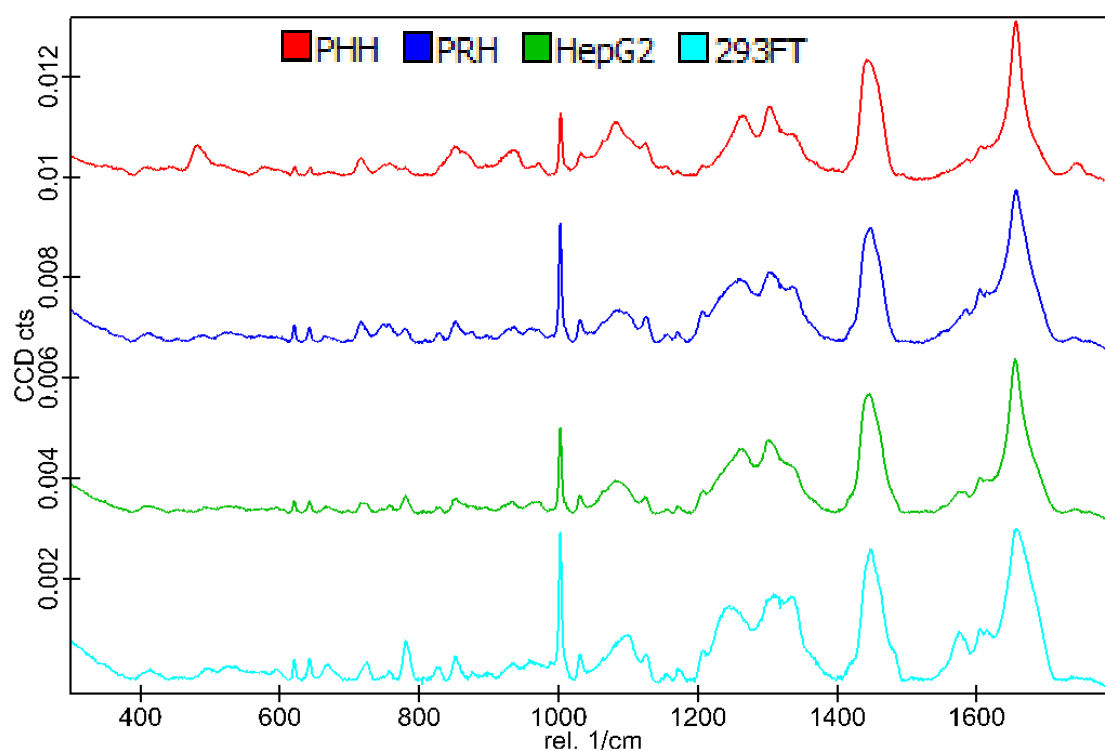


Figure S8. The average Raman spectral of PHH (Lot: 201678901, $n=208$), PRH ($n=201$), HepG2 ($n=204$) and 293FT ($n=205$). (PHH: primary human hepatocytes, PRH: primary rat hepatocytes, HepG2: human liver hepatocellular carcinoma, 293FT: human embryonic kidneys.)

Table S4 Machine learning by stacked (KNN, LDA, PLS, Linear-SVM, RBF-SVM, RF) model to identify cells. Overall accuracy at 81.32% (Lot: 201678901).

	Reference		
	P1	P4	PHH
Model prediction			
P1	106	33	7
P4	38	118	1
PHH	4	2	146
Sensitivity(%)	71.62	77.12	94.81
Specificity(%)	86.97	87.09	98.01